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**Inhibition of CorA dependent Magnesium Homeostasis is cidal in
Mycobacterium tuberculosis.**

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Running Head: Magnesium Homeostasis by *M. tuberculosis* CorA

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30 **ABSTRACT**

31 Mechanisms of magnesium homeostasis in *Mycobacterium tuberculosis* are poorly
32 understood. Herein we describe the characterization of a pyrimidinetrione amide
33 scaffold that disrupts magnesium homeostasis in the pathogen by direct binding to the
34 CorA Mg^{2+}/Co^{2+} transporter. Mutations in domains of CorA that are predicted to regulate
35 the pore opening in response to Mg^{2+} ions conferred resistance to this scaffold. The
36 pyrimidinetrione amides were cidal against the pathogen under both actively replicating
37 as well as non-replicating conditions *in vitro* and were efficacious against the organism
38 during macrophage infection. However, the compound lacked efficacy in infected mice
39 possibly due to limited exposure. Our results indicate that inhibition of Mg^{2+} homeostasis
40 by CorA is an attractive target for tuberculosis drug discovery and encourage
41 identification of improved CorA inhibitors.

42

43 INTRODUCTION

44 Short-course chemotherapy for tuberculosis was adopted by the WHO in 1991 for drug-
45 sensitive disease; requiring 6 months of multi-drug treatment(1). The risk of drug-related
46 toxicity as well as non-compliance increases proportionally over time which has
47 motivated efforts to reduce the treatment duration(2). The current front-line drug
48 regimen consists of drugs that target RNA synthesis (rifampicin), cell wall biosynthesis
49 (isoniazid/ ethambutol) and pyrazinamide (which has an ill-defined mechanism of
50 action) (3, 4). Identification of drugs that target novel pathways in the pathogen has
51 been proposed as a way of developing new drug combinations that could reduce
52 treatment duration(5). To this end, mechanisms that are critical for survival under
53 various growth conditions are predicted to encompass targets that could lead to a rapid
54 decline in bacterial viability *in vivo*.

55 Magnesium is an essential co-factor in a plethora of enzymatic reactions. It
56 serves a structural role in the integrity of nucleic acids and proteins as well as
57 maintaining the integrity of the cell membrane and cell wall(6, 7). In *M. tuberculosis* (*Mtb*)
58 three mutants have been described that exhibited attenuation at low extracellular
59 magnesium concentrations: *perM* encoding a membrane protein of unknown function,
60 deletion of which results in *in vivo* attenuation and inability to grow at low Mg^{2+}
61 concentrations; *phoP* encoding the transcriptional regulator of a two-component system
62 important for complex lipid biosynthesis and virulence; and *mgtC* which encodes a P-
63 type ATPase likely acting as an accessory protein for Mg^{2+} importdeletion of which
64 results in an inability to grow at low Mg^{2+} concentrations at acidic pH and loss of
65 virulence(8-10). The *in vivo* attenuation of these mutants in *Mtb*, as well as other

66 bacterial pathogens with inactivated Mg^{2+} -dependent processes, has been
 67 hypothesized to point to restriction of this essential cation, as a possibly host-mediated
 68 mechanism to control bacterial pathogens(7-10). Indeed, phagosomal Mg^{2+}
 69 concentrations at least during *Salmonella typhimurium* infection have been estimated to
 70 be in the 10-50 micromolar range(11). The predicted low phagosomal Mg^{2+}
 71 concentrations and the increased Mg^{2+} requirement of *Mtb* at low pH, suggests that
 72 access to this nutrient could be growth-limiting *in vivo*. Consequentially inhibitors of
 73 Mg^{2+} homeostasis would affect bacterial survival at every stage of host
 74 pathogenesis(12). The mechanisms by which *Mtb* maintains Mg^{2+} homeostasis are
 75 poorly understood. *Mtb* encodes two Mg^{2+} transporters, CorA and MgtE, neither of
 76 which is apparently essential based on saturating genome transposon mutagenesis
 77 studies(13, 14).

78 In this work, we describe the identification of a compound series that exerts
 79 potent growth inhibition of *Mtb* and inhibits Mg^{2+} uptake via direct binding to the CorA
 80 transporter. A representative of this inhibitor series lacked *in vivo* efficacy against *Mtb* in
 81 mice. This was presumed to be related to insufficient compound exposure *in vivo*.

82

83 RESULTS

84 *Identification of a pyrimidinetrione amide with Gram-positive antibacterial activity*

85 This series, as represented by Compound **1** was identified from a large whole cell
 86 phenotypic screen against *Mtb* and found to have potent antimycobacterial activity on
 87 two distinct growth media (Table 1). The pyrimidinetrione amide core is based on

88 barbituric acid. In order to establish some preliminary structure activity relationship
89 (SAR) analogues were prepared with different amines including a simple aniline (**2**),
90 cyclohexylamine (**3**), butylamine (**4**), benzylamine (**5**), 3-picolylamine (**6**). Except for
91 compound **2**, none showed equivalent antimicrobial activities in both media (Table 1).
92 Although compound **5** showed good potency in GAST medium (a minimal medium
93 lacking BSA), the lack of potency in 7H9 medium suggested high protein binding
94 discouraging further evaluation. The modification of the amide linker to generate either a
95 sulfonylamine (**7**) or a ketone (**8**) also resulted in complete loss of antitubercular activity.

96 Based on the initial SAR described in Table 1, a series of compounds modified in
97 the aniline moiety were further evaluated (Table 2). Interestingly, irrespective of the
98 substituents in the *para*-position of the aniline, compounds **9–14** showed good
99 antitubercular activities. Compounds **15 & 16** which had a substituent in the *ortho*-
100 position also showed activity. While other analogues with heteroaromatic rings instead
101 of a phenyl groups, such as pyridine (**17–19**) and isoxazole (**20**), generally had reduced
102 potency. Because the core is related to barbituric acid, we wanted to rule out non-
103 specific effects due to mitochondrial membrane depolarization(17). We tested for
104 cytotoxicity against HepG2 cells during growth on either glucose or galactose as sole
105 carbon source. Growth on galactose forces cells to use mitochondrial respiration to
106 generate ATP rather than through glycolysis as occurs during growth on glucose(18).
107 This showed that several of the compounds were associated with clear mitochondrial
108 toxicity; evidenced by the enhanced cytotoxicity of compounds during growth on
109 galactose. Cytotoxicity in galactose medium generally tracked with anti-mycobacterial
110 potency (Fig. S1). However, compounds having electron donating alkyl groups, methyl

111 (9) and isopropyl (10), were both selective and had no mitochondrial toxicity based on a
112 lack of inhibition of HepG2 cell growth on either glucose or galactose.,

113 The compound series was not selectively anti-tubercular as seen by its broader-
114 spectrum activity against *Mycobacterium smegmatis*, *Bacillus subtilis* and
115 *Staphylococcus aureus* (Table S1). However, the series lacked activity against Gram-
116 negative bacteria such as *Escherichia coli* and *Pseudomonas aeruginosa* (data not
117 shown), suggesting a mechanism of action unique to Gram positive bacteria.

118

119 *The pyrimidinetrione amide inhibits magnesium uptake in Mtb*

120 In an effort to understand the mechanism of action of the series, resistant mutants were
121 raised to compound 1. Mutants were obtained at a frequency of resistance of 10^{-8} .
122 Three that were confirmed to be more than 100-fold resistant to the compound, were
123 submitted for whole genome sequencing. Two had independent mutations in *corA*
124 (*Rv1239c*), *corA*:E212D and *corA*:A317S, other single nucleotide polymorphisms
125 present showed no evidence of mechanism of action associated resistance (Table 3).
126 The third mutant had an identical SNP profile to the second mutant suggesting that this
127 mutant may have arisen as a sibling.

128 The *corA* gene encodes a putative magnesium and cobalt transporter predicted
129 to be non-essential by genome-wide transposon mutagenesis studies(13, 14). To
130 confirm that SNPs in *corA* were associated with resistance, we raised resistant strains
131 to compounds (1, 10, 12, 13 & 15) and specifically sequenced their *corA* gene. All
132 resistant strains contained point mutations within the *corA* gene (Table S2). To interpret

133 the effect of these mutations on corA function, a MtCorA homology model was
134 generated from the crystal structure of *Thermotoga maritana* CorA (TmCorA) using
135 Phyre2(20). Unexpectedly the mutations did not occur at the divalent cation sensor
136 region where Mg^{2+} ions bind. Instead, the homology model predicted that the mutations
137 occurred within the acidic (E212D) and kink (G299S & M300V/L) regions which are
138 involved in the gating motion of CorA; and the hydrophobic tunnel (A317S) where Mg^{2+}
139 ions flow through (Fig. S2). The mutation data suggested that the pyrimidinetrione
140 amide scaffold affected the gating mechanism of CorA resulting in the inhibition of Mg^{2+}
141 uptake (Fig. S2). This hypothesis was supported by the Mg^{2+} dependence of the growth
142 inhibition in *Mtb* (Fig. 1A) as well as other Gram-positive bacteria (Table S2)

143 Co^{2+} was unable to rescue cells from pyrimidinetrione-dependent growth
144 inhibition (Fig. 1B). The inability of Co^{2+} to rescue growth inhibition was not surprising
145 since there are no known essential cobalt-dependent enzymes in *Mtb*. Magnesium is an
146 essential metal cofactor in many key metabolic enzymes and the specificity of some of
147 the pyrimidinetrione amides for inhibiting bacterial growth and not HepG2 cell growth
148 suggested that magnesium uptake could be a novel target for drug development.

149 To confirm that the series inhibited Mg^{2+} uptake, intracellular concentrations of
150 this cation were measured by inductively-coupled plasma mass spectrometry (ICP-MS).
151 This analysis confirmed that compound **10** treatment generated a concentration-
152 dependent decrease in intracellular magnesium levels (Fig. 1C). Moxifloxacin was
153 chosen as a negative control because it was not expected to affect cell wall or
154 membrane integrity both of which could potentially influence intracellular Mg^{2+} levels.

155 However, moxifloxacin resulted in an unexpected increase in intracellular magnesium
156 concentrations (Fig. 1C) for reasons which were not clear.

157 *The pyrimidinetrione amide- Mg^{2+} complex binds to the CorA transporter*

158 The core of this series shows structural similarity to the pyrimidinedione scaffold
159 associated with Mg^{2+} binding in drugs such as Raltegravir(21). To explore whether this
160 series bound Mg^{2+} ions, we analyzed the UV-Vis spectral response of compound **10**
161 upon titration with Mg^{2+} ions. A clear cation dependent decrease in absorbance at 280
162 nm was seen (Fig. S3). The coordination complex of compound **10** with Mg^{2+} was
163 further investigated by NMR studies; based on the splitting pattern of the signals of the
164 asymmetric pyrimidinetrione moiety, compound **10** forms the enol tautomer when
165 interacting with Mg^{2+} (Fig. S4). The coordination of Mg^{2+} by this scaffold could suggest
166 that the compounds merely act to deplete free Mg^{2+} ions with the *corA* mutations
167 conferring higher affinity for the Mg^{2+} :6H₂O complex. However, arguing against this
168 notion, in the absence of compound the *corA* mutants required a 250-fold higher Mg^{2+}
169 concentration to grow compared to the wild-type strain (Table S3).

170 In order to test whether the Mg^{2+} :pyrimidinetrione amide complex directly bound to
171 CorA, we determined the thermal stability of recombinantly expressed CorA in the
172 presence of Mg^{2+} and compound **10** (Fig. 2). This showed that all of these ligands
173 increased thermal stability of the protein, indicating binding to the transporter (Fig. 2,
174 Table 4). The apparent dissociation constant of CorA with metal ions and compound **10**
175 was determined using fluorescence change based on tryptophan quenching (Fig. S5
176 and Table 5). The results indicated that Co^{2+} was poorly bound to CorA compared to
177 Mg^{2+} whereas compound **10** bound with a 2-fold higher affinity than Mg^{2+} alone,

178 irrespective of the presence of Mg^{2+} . Although this result showed that
179 Mg^{2+} :pyrimidinetrione amide complex bound and stabilized the CorA structure, it was
180 still unclear how this event affected intracellular Mg^{2+} homeostasis, resulting in
181 bactericidal activity. We predicted that mutant strains such as *corA*:E212D could survive
182 exposure to compound **10** as a result of lower binding affinity of the CorA:E212D protein
183 to the Mg^{2+} :compound **10** complex, hereby overcoming the disruption of Mg^{2+}
184 homeostasis. Therefore, we performed similar thermal stability assays and fluorescence
185 shift assays using one of the mutant CorA proteins, the purified recombinant
186 CorA:E212, in the presence of Mg^{2+} , compound **10** or compound **10** with Mg^{2+} . The
187 assay result showed that the Mg^{2+} :pyrimidinetrione amide complex could also bind to
188 the mutated CorA:E212D (Table S4 and S5). This result suggested that the observed
189 mutations on CorA do not have a measurable effect on the binding affinity of the
190 Mg^{2+} :pyrimidinetrione amide complex to this transporter (bearing in mind the limited
191 analytical sensitivity of these assays).

192 *The pyrimidinetrione amide decreases Mtb viability in vitro and ex vivo but lacks*
193 *apparent in vivo efficacy*

194 The pyrimidinetrione amide compound **10** was cidal against actively growing *Mtb* cells
195 showing a 2-log kill over a week of exposure even at concentrations as low as 2-fold the
196 MIC value. Higher concentrations resulted in a more rapid cidal effect with no detectable
197 viable cells remaining as determined by colony counts on solid agar (Fig. 3A). We next
198 sought to interrogate the importance of Mg^{2+} uptake under non-replicating conditions
199 using a non-replicating persistence induced by adaptation to hypoxia with subsequent
200 exposure of cells to compounds under anaerobic conditions. This showed that Mg^{2+}

201 uptake was critical under these conditions as evidenced by the greater than 3-log kill
202 after a week of compound exposure (Fig. 3B).

203 The cidal consequence of disrupting Mg^{2+} homeostasis during active growth as
204 well as non-replicating persistence prompted us to explore their anti-tubercular effect
205 during parasitism of host macrophages. We selected compounds that had no general
206 cytotoxic or mitochondrial toxic effect (compounds **9** and **10**), no general cytotoxic yet
207 with a limited mitochondrial toxicity giving a 10-fold selectivity window (compound **16**) or
208 with limited general cytotoxicity and higher mitochondrial toxicity that still gave us a
209 higher than 10-fold selectivity window (compound **1**) (Table 1). All of these compounds
210 exerted time- and dose-dependent intracellular (*ex vivo*) killing of *Mtb* growing in
211 macrophages (Fig. 3C).

212 The promising activity of compound **10** against *Mtb* during growth in
213 macrophages encouraged us to explore its *in vivo* efficacy. Preliminary pharmacokinetic
214 (PK) analysis indicated the half-life of compound **10** was $2.5 \text{ h} \pm 0.4$ when dosed once
215 daily indicating that at least twice per day (BID) dosing was needed. Unfortunately, this
216 compound was intolerable to mice at high (100 mg/kg) doses. At 30 mg/kg, dosed BID
217 by oral gavage, the compound achieved an AUC of $56 \mu\text{g}\cdot\text{h/mL}$ which gave a AUC/MIC
218 ratio of 250 and plasma concentrations that were almost 10-fold above MIC
219 concentrations at the first trough and just above MIC at 24h ($0.3 \mu\text{g/mL} \pm 0.16 \text{ SD}$; Fig.
220 4A). This dosing schedule was tolerable in naïve mice. Subsequently, *Mtb*-infected mice
221 were treated with 30 or 15 mg/kg BID, however, after multiple BID doses, 30 mg/kg was
222 intolerable to the infected mice necessitating a deescalation to 30 mg/kg once per day
223 for that group for the remaining dosing period. Due to adverse effects, compound **10**

224 could not be administered at higher concentrations and at the highest tolerable
225 concentration, it lacked efficacy after 2 and 4 weeks of treatment in infected mice lungs
226 (Fig. 4B) and spleens (results not shown).

227 DISCUSSION

228 We have identified a pyrimidinetrione amide scaffold with potent *in vitro* activity against
229 *Mtb*. Resistant mutants mapped to the CorA transporter with Mg^{2+} supplementation
230 abrogating the growth inhibition in a concentration-dependent fashion. Using
231 thermostability assays and fluorescence shift assays, we were able to demonstrate that
232 this inhibitor binds to CorA in the presence and absence of Mg^{2+} . The ligand binding site
233 for this inhibitor remains to be identified but we were also able to show that resistance-
234 conferring amino acid mutations had no effect on this ligand binding. Instead, the
235 mutations mapped to the acidic and kink regions of the cytoplasmic domain of the CorA
236 transporter which are involved in the gating motion of the pore, indicating that mode of
237 resistance were not due to changes in CorA-ligand interaction, but likely as a result of
238 perturbation of the gating motion in response to Mg^{2+} dissociation from the protein. Low
239 cytosolic Mg^{2+} concentrations result in cation dissociation with resultant channel
240 opening driven by this gating mechanism. In line with a loss of regulation of the pore
241 opening of the cation channel, the *corA*:E212D and *corA*:A317 mutants required a 250-
242 fold higher Mg^{2+} concentration in the medium than parental wild-type cells for growth
243 (results not shown). During preparation of this manuscript, a report was published that
244 found similar mutations in CorA to a pyrimidinetrione as well as 4-hydroxyquinoline
245 scaffold(19). These mutations were predicted to result in channel opening thereby
246 resulting in increased cytosolic Mg^{2+} concentrations. Similar to our work, cytosolic

247 concentrations of both free and bound Mg^{2+} were decreased complementing our data
248 that these inhibitors directly influence the influx/efflux of this cation.

249 The observation that the metabolic consequence of CorA inhibition is both growth arrest
250 and death was surprising since there is an alternative Mg^{2+} transporter encoded by
251 *mgtE* in the bacterium. The finding that only mutations in *corA* could confer resistance
252 suggests that regulation of Mg^{2+} homeostasis by this cation is predominantly conferred
253 by the CorA transporter where changes in MgtE levels do not occur rapidly enough to
254 overcome this dysregulation of Mg^{2+} homeostasis. In contrast, during genetic
255 inactivation of *corA*, compensatory genetic changes could occur to alter levels of MgtE
256 as cells are gradually depleted of CorA protein. The finding that *corA* but not *mgtC* or
257 *mgtE* is conserved in *Mycobacterium leprae*, a mycobacterium with a minimal genome,
258 points to the conserved nature of Mg^{2+} homeostasis by this transporter. Despite this,
259 there is no evidence for the *in vivo* essentiality of *Mtb corA* during mouse
260 pathogenesis(22). Intriguingly, these compounds were only active against Gram positive
261 bacteria tested and not Gram negatives. CorA is widely distributed in bacteria including
262 *E. coli* and *P. aeruginosa*(6). Lack of activity against these Gram negative microbes
263 could be due to lack of binding to their CorA counterpart, different mechanisms of Mg^{2+}
264 homeostasis, lack of transport of the pyrimidinetrione amides across the outer
265 membrane or their efflux by multidrug efflux pumps.

266 Our results emphasize the importance of Mg^{2+} homeostasis during active
267 replication as well as non-replicating persistence *in vitro* and also during parasitism of
268 host macrophages. Free Mg^{2+} concentrations in the lung were determined to be $0.67 \pm$
269 0.08 mM (data not shown), a value similar to reported concentrations in plasma(23),

270 which is below the level that overcomes the growth inhibition by the pyrimidinetrione
271 amides. The lack of *in vivo* efficacy was disappointing although the pharmacokinetic
272 studies suggested that blood serum concentrations were only sufficiently above the MIC
273 for half of the dosing period when extrapolated from the PK parameters (Fig. 4A). The
274 concentration of compound in the cellular lesions may further be well below the MIC
275 level during this dosing period. In addition, despite the lack of cytotoxicity of the
276 compound, clinical signs of toxicity were evident when mice were dosed at 100 mg/kg
277 (results not shown) prohibiting additional studies to probe the *in vivo* efficacy of this
278 compound.

279 In conclusion, our results demonstrate the critical role of CorA in maintaining
280 Mg^{2+} homeostasis and its attractiveness as a drug target. Despite the *in vitro* and *ex*
281 *vivo* efficacy of the pyrimidinetrione amides, our data suggests that an alternative
282 scaffold that inhibits CorA function would need to be identified in order to perform the
283 proof of concept experiments that establish the validity of this target *in vivo*.

284

285 MATERIALS AND METHODS

286 *Chemistry.* All chemicals and reagents were purchased from Aldrich (Sigma Aldrich, St.
287 Louis, MO, USA), Lancaster (Alfa Aesar, Johnson Matthey Company, Ward Hill, MA,
288 USA) and were used without further purification. Commercially available analogues of
289 pyrimidinetrione amide were purchased from Enamine LLC (NJ, USA), Bionet America,
290 Inc. (CA, USA), and MicroCombiChem GmbH (Wiesbaden, Germany). Reactions were
291 monitored by TLC, performed on silica gel glass plates containing 60 GF-254, and

292 visualization on TLC was achieved by UV light or iodine indicator. Column
293 chromatography was performed with Merck (60-120) mesh silica gel. ^1H NMR spectra
294 were recorded on Varian Mercury-300 NMR Spectrometer and chemical shifts are
295 reported in ppm downfield from internal TMS standard. Routine mass analyses (LRMS)
296 were performed on HP Agilent LC/MS series 1100 system equipped with a reverse
297 phase column (Agilent Poroshell 120 EC-C18, 2.7 μm , 50 \times 2.1 mm) and photodiode
298 array detector using electrospray ionization (ESI). All the reported compounds were
299 pure and > 95%.

300 *Synthesis of compound 12, 14–20(24)*

301 To a solution of 1,3-dimethylpyrimidine-2,4,6(1*H*,3*H*,5*H*)-trione (1.0 eq) in DCM (0.1 M)
302 was added pyridine (1.2 eq) and 4-dimethylaminopyridine (0.1 eq) at 0°C.
303 Ethylcarbonochloridate (1.0 eq) was added dropwise to the reaction mixture at the same
304 temperature over 1h. After stirring for 12 h at 0°C, the reaction mixture was allowed to
305 warm to room temperature and then diluted with DCM and washed with water. The
306 organic layer was dried over Na_2SO_4 and evaporated *in vacuo*. The crude mixture was
307 recrystallized from ethanol to get ethyl 1,3-dimethyl-2,4,6-trioxohexahydropyrimidine-5-
308 carboxylate as a white solid (1.09 g, 75%). ^1H NMR ($\text{DMSO}-d_6$, 300 MHz) δ 4.28 (qt, J =
309 6.9 Hz, 2H), 3.16 (s, 3H), 1.27–1.23 (t, J = 6.9 Hz, 3H); LCMS (ESI) m/z 229 $[\text{M}+\text{H}]^+$.

310 To a solution of ethyl 1,3-dimethyl-2,4,6-trioxohexahydropyrimidine-5-carboxylate (1.0
311 eq) in toluene (0.5 M) was added each amine (1.0 eq) and TEA (1.2 eq). The reaction
312 mixture was refluxed for about 12 h. After cooling down to room temperature, the
313 reaction mixture was diluted with ethyl acetate and then washed with water. The organic

layer was dried over Na₂SO₄ and evaporated *in vacuo*. The crude mixture was recrystallized from acetonitrile to get the desired products, **12**, **14–20**.

1,3-Dimethyl-2,4,6-trioxo-N-(4-phenoxyphenyl)hexahydropyrimidine-5carboxamide (12).

White solid (75%): ¹HNMR (CDCl₃, 300 MHz) δ 11.92 (s, 1H), 7.46 (d, *J* = 6.9 Hz, 2H), 7.35 (t, *J* = 7.2 Hz, 2H), 7.12 (t, *J* = 7.5 Hz, 1H), 7.03–7.00 (m, 4H), 3.41 (s, 3H), 3.38 (s, 3H); LCMS (ESI) *m/z* 368 [M+H]⁺.

N-(4-acetylphenyl)-1,3-dimethyl-2,4,6trioxohexahydropyrimidine-5-carboxamide (14).

White solid (79%): ¹HNMR (CDCl₃, 300 MHz) δ 12.10 (s, 1H), 7.97 (d, *J* = 8.7 Hz, 2H), 7.63 (d, *J* = 8.7 Hz, 2H), 3.41 (s, 3H), 3.37 (s, 3H), 2.59 (s, 3H); LCMS (ESI) *m/z* 318 [M+H]⁺.

N-(4-bromo-2-fluorophenyl)-1,3-dimethyl-2,4,6-trioxohexahydropyrimidine-5-

carboxamide (15). White solid (46%): ¹HNMR (CDCl₃, 300 MHz) δ 12.04 (s, 1H), 7.98 (t, *J* = 8.5 Hz, 1H), 7.38–7.26 (m, 2H), 3.42 (s, 3H), 3.38 (s, 3H); LC-MS: LCMS (ESI) *m/z* 370 [M-H]⁻.

1,3-Dimethyl-2,4,6-trioxo-N-(2,4,6-trichlorophenyl)hexahydropyrimidine-5-carboxamide

(16). White solid (48%): ¹HNMR (CDCl₃, 300 MHz) δ 11.35 (s, 1H), 7.44 (s, 2H), 3.40 (s, 3H), 3.39 (s, 3H); LCMS (ESI) *m/z* 378 [M+H]⁺.

N-(6-chloropyridin-3-yl)-1,3-dimethyl-2,4,6trioxohexahydropyrimidine-5-carboxamide

(17). White solid (88%): ¹HNMR (CDCl₃, 300 MHz) δ 11.94 (s, 1H), 8.52 (d, , *J* = 2.4 Hz, 1H) 8.01–7.96 (dd, *J* = 2.7 and 8.7 Hz , 1H), 7.35 (d, *J* = 8.4 Hz, 1H), 3.41 (s, 3H), 3.37 (s, 3H); LCMS (ESI) *m/z* 311 [M+H]⁺.

335 1,3-dimethyl-N-(6-methylpyridin-2-yl)-2,4,6-trioxohexahydropyrimidine-5-carboxamide
 336 (18). White solid (71%): ¹HNMR (CDCl₃, 300 MHz) δ 12.13 (s, 1H), 7.70 (d, *J* = 7 Hz,
 337 1H), 7.63 (t, *J* = 7.8 Hz, 1H), 6.98 (d, *J* = 7.5 Hz, 1H), 3.39 (s, 6H), 2.50 (s, 3H); LCMS
 338 (ESI) *m/z* 291 [M+H]⁺.

339 1,3-Dimethyl-2,4,6-trioxo-N-(4(trifluoromethyl)pyridin-2-yl)hexahydropyrimidine-5-
 340 carboxamide (19). White solid (73%): ¹HNMR (CDCl₃, 300 MHz) δ 12.18 (s, 1H), 8.55
 341 (d, *J* = 5.1 Hz, 1H), 8.27 (s, 1H), 7.32 (d, *J* = 4.8 Hz, 1H), 3.43 (s, 3H), 3.39 (s, 3H);
 342 LCMS (ESI) *m/z* 345 [M+H]⁺.

343 1,3-Dimethyl-N-(5-methylisoxazol-3-yl)-2,4,6-trioxohexahydropyrimidine-5-carboxamide
 344 (20). White solid (69%): ¹HNMR (CDCl₃, 300 MHz) δ 12.04 (s, 1H), 6.52 (s, 1H), 3.41 (s,
 345 3H), 3.36 (s, 3H), 2.43 (s, 3H); ¹³C NMR (CDCl₃, 75 MHz) δ 169.1, 167.8, 162.8, 161.7,
 346 158.4, 91.7, 81.6, 28.5, 12.1; LCMS (ESI) *m/z* 281 [M+H]⁺.

347 *Strains, media, and materials.* *Mtb* H37Rv (ATCC 27294) was used for all experiments.
 348 To determine the antimicrobial spectrum of the compounds, *B. subtilis* and *S. aureus*
 349 was used as gram positive strains, *E. coli* and *P. aeruginosa* were used as gram
 350 negative strains. For liquid cultures, *Mtb* was grown in 7H9 medium which consists of
 351 Middlebrook 7H9 (Becton Dickinson) broth base supplemented with ADC [albumin
 352 (50 g/L)/dextrose (20 g/L)/NaCl (8.1 g/L)], 0.2% glycerol] and 0.05% Tween 80. Solid
 353 agar media (7H11/OADC) for *Mtb* growth consisted of Middlebrook 7H11 (Becton
 354 Dickinson) agar base supplemented with OADC [ADC with 0.06% oleic acid]. GAST
 355 (glycerol-alanine-salts-Tween 80) was used as defined minimal liquid growth media for
 356 *Mtb*(25). Mg²⁺-free GAST media made by omitting MgCl₂·6H₂O from the medium

357 composition. The low pH nitrosative stress medium contained bovine serum albumin
358 fraction V (5 g/L)/butyric acid (2.5 mM)/NaNO₂ (6.9 mg/mL)/NaCl (0.81 g/L)/0.05%
359 Tyloxapol, pH adjusted to 6.0. MIC determination was performed as previously
360 described(25). Luria-Bertani (LB) broth (Difco, New Jersey, USA) was used for MIC of
361 Gram positive and negative strains. The screening conditions under which the original
362 pyrimidinetrione hits were identified have been previously described but in brief
363 consisted of 7H9 medium with readout of *Mtb* growth after 3 days of compound
364 exposure(26).

365 *Generation and characterization of pyrimidinetrione resistant mutants.* To generate
366 spontaneously resistant mutants to the compounds, 50 ml of *Mtb* was grown to OD_{650nm}
367 0.2. Harvested cells were resuspended in 500 µL of media and 100 µL aliquots (10⁹
368 cells) plated on 7H11/OADC containing compound **1** at concentrations corresponding to
369 5x, 10x, 20x MIC. Appropriate dilutions of the cell suspensions were also plated on
370 drug-free plates to calculate bacterial numbers in the suspensions for determination of
371 frequencies (resistance frequency being the number of resistant mutants obtained per
372 number of bacteria plated). Agar plates were incubated at 37°C for 4 weeks. After 4
373 weeks, 4 colonies growing on 7H11/OADC containing 10x MIC compound **1** were
374 inoculated in 7H9 media and resistance confirmed by MIC determination. Genomic DNA
375 of mutants was isolated by the CTAB method(27). Whole genome sequencing was
376 performed and analyzed as described(28).

377 *Cytotoxicity in HepG2 cells.* Cytotoxicity in HepG2 cells was measured by CellTiter-
378 Glo™ Luminescent Cell Viability Assay (Promega). For glucose media, 20,000 HepG2
379 cells were seeded in 96 well white flat-bottom plates (Corning incorporated) in DMEM

380 GlutaMAX (Gibco) supplemented with 10% fetal bovine serum, 20 mM HEPES, and 0.5
381 mM sodium pyruvate. For galactose media, HepG2 cells were pre-cultured in galactose
382 DMEM and seeded in same plate with glucose media. Galactose DMEM consists of
383 glucose-free DMEM (Gibco) supplemented with 10% fetal bovine serum, 10 mM D-
384 galactose, 2 mM L-glutamine, 5 mM HEPES, and 0.5 mM sodium pyruvate. The plates
385 were incubated at 37°C with 5% CO₂. The following day, 2-fold serial dilutions of
386 compound up to 100 µM were added in the respective growth medium in duplicate and
387 cells incubated for 24 hours. After this, 20 µL of CellTiter-glo reagent was added to each
388 well followed by incubation at 37°C with 5% CO₂ for 10 min. Cytotoxicity was
389 determined by measuring luminescence using Fluostar Optima FL (BMG Labtech).

390 *In Silico Analysis.* Amino acid sequence of MtCorA (Rv1239c) was retrieved from
391 Mycobrowser database (<https://mycobrowser.epfl.ch/>). FASTA format of MtCorA protein
392 sequence was submitted to PHYRE2 protein fold recognition server. Ninty nine
393 sequence alignment hits were obtained. The sequence of TmCorA was selected as a
394 single highest scoring template and 332 residues (91% of MtCorA sequence) were
395 modelled with 100% confidence by the structure of TmCorA.

396 *Magnesium and cobalt concentration dependent MIC.* We used Mg²⁺-free GAST media
397 to test the magnesium-dependence of the MIC of the compounds. Two-fold
398 concentrated GAST generated without the addition of magnesium chloride hexahydrate
399 was prepared. An equal volume of magnesium chloride in sterile distilled water was
400 added to the GAST medium to give final concentrations ranging from 0.12 mM up to
401 240 mM. MIC determinations for the compounds were set up in duplicate as previously
402 described for each MgCl₂ concentration(29). Briefly, an equal volume of medium (50 µL

403 GAST at the defined MgCl_2 concentration) with compound serially diluted in duplicate
404 from 50 to 0.024 μM was added to wells of a round-bottom 96-well plate (Nunc). *Mtb*
405 culture in GAST at $\text{OD}_{650\text{nm}}$ of 0.2 was diluted 1: 500 in 2x Mg^{2+} -free GAST and 50 μL
406 added to each well of the 96 well plates. Plates were incubated under 5% of CO_2 at
407 37°C . Growth was measured using an inverted enlarging mirror at 1, 2 and 3 weeks with
408 MIC taken as the concentration that completely inhibited all visible growth. For cobalt
409 dependent MIC test, CoCl_2 was added in GAST media from 0.001 to 5 mM
410 concentration and MIC performed as above.

411 *Magnesium concentration measurement by inductively coupled plasma - mass*
412 *spectroscopy (ICP-MS).* We measured magnesium concentrations in *Mtb* cells treated
413 by inductively coupled plasma - mass spectroscopy (ICP-MS). Two liters of *Mtb* was
414 grown to $\text{OD}_{650\text{nm}}$ 0.6 in 7H9. Five hundred milliliter of culture volumes were incubated
415 with 1x, 10x MIC of compound **10**, 10x MIC of moxifloxacin or an equivalent volume of
416 DMSO vehicle control (0.1% v/v) for 24 hours in 2 L capacity rolling bottles. Cells were
417 harvested and washed three times with 1/10 volume phosphate buffered saline
418 containing 0.05% Tween 80 (PBST). The wet weight of *Mtb* cell pellets was recorded to
419 normalize magnesium concentration prior to autoclaving. ICP-MS analysis was serviced
420 by Elementary Analysis Inc. (Lexington, KY, USA)

421 *UV spectrum scanning to test magnesium binding of pyrimidinetrione.* One milliliter of
422 100 μM **10** dissolved in 100 mM HEPES buffer was filled in 1 cm path length quartz
423 crystal cuvettes for UV scanning. The absorbance of the compound **10** solution was
424 scanned from 200 nm to 400 nm wavelengths by a Varian Cary 300 Bio UV/VIS
425 spectrophotometer. We sequentially added 10 μL of 1 mM MgSO_4 to the compound

426 solution to observe changes in absorbance spectrum. A standard curve was plotted with
427 $A_{280\text{nm}}$ of 5, 10, 50 and 100 μM of compound **10**. Absorbance at 280 nm was used to
428 calculate compound **10** after correcting for changes in volume and plotted in a graph.

429 *Minimum magnesium requirement test.* *Mtb* H37Rv, *corA*:E21D, and *corA*:A317S
430 strains were cultured in GAST to $\text{OD}_{650\text{nm}}$ 0.2 at 37°C. Fifty microliter of Mg^{2+} -free
431 GAST was distributed in every well of a round-bottom 96-well plate (Nunc). Fifty μL
432 of 240 mM of MgCl_2 dissolved in Mg^{2+} -free GAST was added to the wells of the first
433 column of this plate and 50 μL volumes transferred from column 1 to column 12
434 generating a 2-fold serial dilution series of MgCl_2 across the plate. *Mtb* cultures were
435 diluted 1: 500 in Mg^{2+} -free GAST and 50 μL added to each well of the 96 well plates.
436 Growth was measured using an inverted enlarging mirror at 1, 2 and 3 weeks with the
437 minimum Mg^{2+} concentration to support growth taken as the lowest concentration that
438 supported evidence of any visible growth.

439 *Cloning, expression and purification of MtCorA.* The *M. tuberculosis corA* gene,
440 *rv1239c*, was cloned into the pET28a vector. The gene was amplified by PCR using
441 Herculase II Fusion DNA polymerase (Agilent) and the following primers: forward-5'-
442 CGCGGCAGCCATATGTTCCCAGGGTTT-3' and reverse-5'-
443 TGCGGCCGCAAGCTTCTAGAGCCAGTTTCT-3'. The PCR-amplified gene was
444 inserted between two restriction sites, *NdeI* and *HindIII*, using In-Fusion® HD Cloning
445 Plus (Takara). Site-directed mutagenesis to obtain gene encoding *corA*:E212D mutant
446 was performed. MtCorA contained the N-terminal poly-histidine tag. The plasmids
447 carrying the wildtype and mutant *corA* gene were transformed into in OverExpress™
448 C41(DE3) (Lucigen). The cells were grown in LB media at 37°C. The induction was

449 initiated with 1mM IPTG at OD_{600nm} 0.7-0.8 and after 6 h, the cells were harvested by
450 centrifugation at 3500 rpm for 20 min. The cell pellet was resuspended in lysis buffer
451 containing 50 mM Tris-HCl pH 8.0, 150 mM NaCl, 0.5 mM DTT and SIGMAFAST™
452 Protease Inhibitor Cocktail Tablets, EDTA-Free (Sigma-Aldrich). After sonication, the
453 lysed cells were pelleted by centrifugation at 20,000 g for 60 min and the membrane
454 was isolated from the supernatant by ultracentrifugation at 142,000 g for 60 min. The
455 isolated membrane was then solubilized in the resuspension buffer, 50 mM Tris-HCl pH
456 8.0, 300 mM NaCl, 20 mM imidazole, 0.5 mM DTT, with 1% (w/v) dodecyl maltoside
457 (DDM) (Anatrace). The resulting supernatant was loaded onto a Ni-NTA column. The
458 column was washed with wash buffer consisting of 50 mM Tris-HCl pH 8.0, 300 mM
459 NaCl, 40 mM imidazole, 0.5 mM DTT, with 0.1% (w/v) DDM. The CorA or CorA mutant
460 protein was eluted with elution buffer (50 mM Tris-HCl pH 8.0, 300 mM NaCl, 300 mM
461 imidazole, 0.5 mM DTT, with 0.05% (w/v) DDM) and the resulting supernatant fractions
462 containing CorA were identified using SDS-PAGE, pooled and dialyzed against dialysis
463 buffer (50 mM Tris-HCl pH 8.0, 100 mM NaCl, 0.5 mM DTT, with 0.05% (w/v) DDM). The
464 protein was finally purified on a Superdex 200 16/60 (GE Healthcare) in 50 mM Tris-HCl
465 pH 8.0, 300 mM NaCl, 0.5 mM DTT, with 0.05% (w/v) DDM.

466 *Thermostability test of CorA with pyrimidinetriones.* Purified MtCorA samples were
467 diluted to a concentration of 0.5 mg/mL. The solution was aliquoted into 30 µl volumes,
468 with 1 µL of a stock solution of different Mg²⁺ concentrations added to the aliquot.
469 Samples were incubated for 20 min at room temperature and then further incubated for
470 10 min at every increment of 5°C up to 95°C in a thermocycler. The samples were

471 transferred to a 96-well filter plate (0.45 µm) (Millipore) to remove precipitated proteins.

472 The yield of the filtered protein was analyzed by SDS/PAGE.

473 *Substrate Binding Measured by Fluorescence Quenching.* The dissociation constants
474 between CorA and its ligands, Mg²⁺, Co²⁺ and compound **10**, were determined by the
475 ligand dose-dependent fluorescence quenching assay (Cary Varian Eclipse
476 fluorescence spectrophotometer). CorA was titrated with increasing concentrations of
477 ligands and fluorescence emission was measured at ~350 nm following excitation at
478 283 nm (bandwidths of 5nm). The fluorescence quenching ΔF versus ligands
479 concentration [S] was plotted and K_d(app) of CorA to ligands were calculated by fitting
480 the data to an equation,

$$\Delta F = (F_0 - F) = \frac{\Delta F_{\max} * [S]}{K_d(\text{app}) + [S]}$$

481 where F₀ and F are the fluorescence intensities in the absence and presence of
482 substrates, respectively, [S] is the ligand concentration, and K_d(app) is the apparent
483 dissociation constant of CorA to ligands.

484 *Efficacy study in vitro, ex vivo and in vivo.* *In vitro* efficacy of compounds against *Mtb*
485 was measured during aerobic growth as well as during anaerobic non-replicating
486 persistence. Time kill kinetics were measured under aerobic conditions as follows.
487 Logarithmically growing *Mtb* (OD_{650nm} 0.2) was diluted 1,000-fold in 7H9 media and 1 ml
488 volumes exposed to 1x, 2x, 5x, 10x, and 20 x MIC compound **10** up to 7 days in
489 duplicates. Samples were collected and plated on 7H11/OADC plates for CFU
490 enumeration at 1, 2, 4 and 7 days of treatment. For anaerobic conditions, *Mtb* was
491 cultured in the self-generated oxygen-depletion model as previously described(29).

492 One milliliter volumes of anaerobic *Mtb* culture was exposed 1 week to 1x, 5x, 10x, 20x
493 and 50x MIC compounds **9**, **10**, and **20** in an anaerobic chamber (Microbiology
494 International) followed by CFU enumeration as above.

495 For *ex vivo* efficacy testing, J774 cells (6.6×10^4 cells/well) were seeded in flat-bottom
496 24 well plates (Corning incorporated) in J774 growth medium consisting of DMEM
497 GlutaMAX (Gibco) supplemented with 10% fetal bovine serum, 20 mM HEPES + 0.5
498 mM sodium pyruvate. *Mtb* H37Rv with confirmed high levels of phthiocerol
499 dimycocerosate was grown in 7H9 to an OD_{650nm} of 0.5, filtered through a 5 μ m sterile
500 filter to ensure a single cell suspension, cell density confirmed at OD_{650nm}, and diluted in
501 J774 growth medium to 4.4×10^5 CFU/mL. Aliquots of 0.1 mL of the *Mtb* suspension
502 was added to each well of the 24-well plate giving an MOI 1:1.5. After overnight
503 incubation, medium was aspirated and monolayers were washed twice in pre-warmed
504 PBS (pH 7.4) twice. At this point, four wells were used for CFU enumeration as
505 described below. To the remaining wells, 1mL J774 growth medium containing
506 compound or vehicle (DMSO) control was added to each well. Each compound was
507 tested at 1, 5, and 10-fold its MIC concentration in duplicate wells for each time point
508 and concentration. Rifampicin (0.5 and 5 μ g/mL) was used as positive control. Cells
509 were incubated at 37°C, 95% humidity, 5% CO₂ incubator for 3 and 7 days. Media was
510 replenished at days 3 and 5 taking care not to remove any macrophages since the
511 pyrimidinetrione amides resulted in loss of adherence of macrophages over time. After 3
512 or 7 days of incubation, medium was removed taking care not remove macrophages and
513 1mL of 7H9 medium containing 0.1% SDS added to each well to ensure macrophage
514 lysis. After 5 minutes, lysate was rapidly mixed to shear eukaryotic DNA, diluted in 7H9

515 and plated on 7H11/OADC plates. Colony counts were enumerated after 4 weeks of
516 incubation at 37°C.

517 Mouse studies were carried out in accordance with the Guide for the Care and Use of
518 Laboratory Animals of the National Institutes of Health under Animal study protocol
519 numbers LCIM 4E. Pharmacokinetic analyses on 20 C57BL/6J mice were done by
520 administering 30 mg/kg compound **10** suspended in 10% (2-Hydroxypropyl)- β -
521 cyclodextrin with 50 mM tricine (pH 8.2) twice per day (with a separation of 7 hours) by
522 oral gavage followed by blood sampling at 0.5, 1, 2, 4, 8, 12 and 24 hours. 0.1 mL of
523 blood was collected twice from each animal over the experiment into Li-Heparin
524 Microvette tubes (Sardtedt AG & Co; Numbrecht, Germany) and centrifuged to prepare
525 to plasma. Tolerability of compound **10** was tested by dosing 5 naïve mice twice per
526 day with 100 or 30 mg/kg for 7 days with observation for 7 subsequent days. For
527 evaluation of *in vivo* efficacy, C57BL/6J mice were infected by the aerosol route as
528 previously described(30). After 14 days, groups of 10 mice were dosed with compound
529 **10** given by oral gavage of either 15 mg/kg or 30 mg/kg twice a day. The higher dose
530 group developed signs of distress after 1 week and so that group was dose deescalated
531 to 30 mg/kg once a day. Control groups were dosed with vehicle control (10% (2-
532 Hydroxypropyl)- β -cyclodextrin) with 50 mM tricine (pH 8.2) or 10 mg/kg Rifampicin. After
533 2 and 4 weeks of treatment, groups of 5 mice were euthanized and appropriate dilutions
534 in 7H9 medium of lung and spleen homogenates plated on 7H11/OADC plates for CFU
535 enumeration.

536 *Pharmacokinetic analysis of pyrimidinetriones.* Pharmacokinetic plasma samples were
537 analyzed by an Agilent 1200 Infinity HPLC with Agilent 6460C Triple quadrupole mass

selective detector (LC-QqQ) utilizing electrospray ionization (ESI). Samples were prepared by mixing 50 μ L serum with 50 μ L labetalolTM internal standard (IS), 20 μ L water or spike solution, followed by 400 μ L acetonitrile/methanol 3:1 for protein precipitation. Samples were centrifuged 13 k rpm x 5 min and 5 μ L supernatant injected into a 2.1 x 50 mm EclipsePlus C18 1.8 μ m column utilizing aq. 0.1% formic acid (solvent A) and acetonitrile w 0.1% formic acid (solvent B). Solvent gradient program was 8% B hold 0.5 minutes to 95% B in 4.5 minutes with flow rate 0.8 ml/min.

MRM transition for compound **10** was detected as [M-H]⁻ precursor (Q1), CEV 24V, and 155 product ion(Q2) negative mode, and IS detected as M+H⁺ precursor (Q1), CEV 8V, and 311.2 product ion (Q2) positive mode. PK parameters were calculated using PK macros in Excel.

549

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558 ABBREVIATIONS

559 *Mtb*, *Mycobacterium tuberculosis*; Kan, kanamycin; INH, isoniazid; RIF, rifampicin; MIC,
560 minimum inhibitory concentration; AUC, area under the curve; CFU, colony-forming
561 units; ICP-MS, inductively-coupled plasma mass spectrometry.

562

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653
654

655 **TABLE 1** Antitubercular activity of pyrimidinetrione amide analogs

Compound ^a	Chemical Structure	MIC (μM)	
	R	7H9 ^b	GAST ^c
1		0.2	0.29
2		2.24	1.49
3		>50	9.4
4		25	12.5
5		50	0.39
6		>50	>50
7		>50	37
8		>50	>50

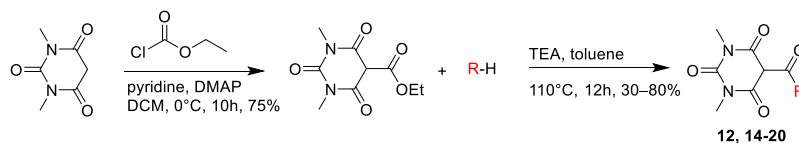
656 ^a All compound **1–8** are known molecules. MIC of compound against *Mtb* H37Rv in ^b Middlebrook 7H9/BSA
 657 containing glucose/glycerol/Tween80, ^c GAS medium with Tween80.

658

659

660 **TABLE 2** Antitubercular activity and cytotoxicity of pyrimidinetrione amide analogs

661



Compound ^a	Chemical Structure	MIC (μM)				Cytotoxicity (IC ₅₀ , μM)	
	R	7H9 ^b	GAST ^c	Butyrate ^d	7H9 (<i>corA</i> :E212D) ^e	Glucose ^f	Galactose ^g
1		0.20	0.29	0.10	>50	29.5	5.50
2		2.24	1.49	ND	ND	ND	ND
9		1.20	0.78	1.56	>50	>100	>100
10		0.59	0.20	3.13	6.25	>100	>100
11		3.10	1.56	3.13	3.10	72.5	23.0
12		0.78	0.20	0.59	6.25	26.6	3.50
13		0.20	0.07	1.17	3.13	24.6	3.80
14		3.10	4.69	3.13	25.0	8.54	2.10
15		0.78	0.20	1.17	9.38	19.0	5.20
16		6.30	1.17	6.25	>50	>100	50
17		4.70	3.13	3.13	>50	>100	28.0
18		>50	>50	>50	>50	ND	ND
19		>50	>50	>50	>50	ND	ND
20		9.38	2.34	50	>50	ND	ND

662 ^a Compound **1**, **2**, **9–11**, **13** are known molecules. MIC of compounds against *Mtb* H37Rv in ^bMiddlebrook 7H9/BSA
 663 containing glucose/glycerol/Tween80, ^cGAS medium with Tween80, ^dMiddlebrook 7H9/BSA/Tyloxapol/butyrate/0.1
 664 mM nitrite/pH 6.0. ^e MIC of compounds against *CorA*:E212D mutant *Mtb* strain in Middlebrook 7H9/BSA containing

665 glucose/glycerol/Tween80. ^f Cytotoxicity of compound tested against HepG2 cells in DMEM/10% FBS supplemented
 666 with glucose. ^g Cytotoxicity of compound tested against HepG2 cells in DMEM/10% FBS with galactose.

667

668 **TABLE 3** Mutations identified in pyrimidinetrione amide-resistant *Mtb* strains

Strain	Polymorphisms
E2.1	<i>corA</i> :E212D, 4221182:T>C (-30 bp upstream of <i>Rv3776</i>)
E2.2	<i>corA</i> :A317S, <i>Rv0275c</i> :S233S, 3281706 -G (in aa 342/2111 of <i>mas</i>)
E2.6	Same as E2.2

669

670 **TABLE 4** Thermostability of CorA in the presence of metal cations, EDTA and compound **10**.

CorA + additive	Melting temperature (°C)
none	55
0.125 mM MgCl ₂	70
1 mM compound 10	≥95
1 mM compound 10 / 0.125 mM MgCl ₂	≥95
1 mM EDTA / 0.125 mM MgCl ₂	≥95

671

672 **TABLE 5** The apparent dissociation constant ($K_d(\text{app})$) of CorA with metal cations and
 673 compound **10**.

CorA additives in fluorescence shift assay	$K_d(\text{app})$ μM
Mg ²⁺	47.5 \pm 3.0
Co ²⁺	205 \pm 18
Compound 10	20.0 \pm 4.3
Compound 10 / 1 mM Mg ²⁺	24.1 \pm 6.3
Compound 10 / 1 mM Co ²⁺	21.8 \pm 4.1

674

675

676 **FIGURE LEGENDS**

677 **FIG 1** The pyrimidinetrione amide scaffold inhibits Mg^{2+} uptake by *Mtb*. (A) Magnesium
678 dependence of the pyrimidinetrione amide MIC against *Mtb* performed in Mg^{2+} -free
679 media with the indicated concentrations of Mg^{2+} . (B) Co^{2+} dependence of the
680 pyrimidinetrione amide MIC against *Mtb* performed in Co^{2+} -free media with the indicated
681 concentrations of Co^{2+} . (C) Fold change in intracellular Mg^{2+} concentration as measured
682 by ICP-MS compared to vehicle (DMSO) control after 24h of treatment with moxifloxacin
683 (Mox) or compound **10** at 1- or 10-fold MIC concentration in 7H9 medium which
684 contains 0.4 mM Mg^{2+} . Significance between groups was determined by Tukey's
685 Multiple Comparison Test with significance set at P values less than 0.05.

686 **FIG 2** Magnesium and the pyrimidinetrione scaffold bind to the CorA transporter
687 protein. The thermostability of recombinantly expressed CorA from *Mtb* was measured
688 with (A) no ligand, (B) 0.125 mM $MgCl_2$ and (C) 0.125 mM $MgCl_2$ with or without 1.0 mM
689 compound **10**. Protein was pre-incubated at room temperature with the indicated
690 ligands before heating to the indicated temperatures in a thermocycler. Protein
691 precipitates were removed by filtration and soluble protein analyzed by SDS-PAGE.

692 **FIG 3** The pyrimidinetrione amide scaffold is cidal against replicating as well as non-
693 replicating *Mtb* *in vitro* and *ex vivo*. (A) Kill kinetics of compound **10** against *Mtb* during
694 active replication *in vitro* as compared to negative (DMSO) and positive (5 μ g/mL RIF)
695 controls. (B) Anaerobic cidal activity of compounds **9**, **10** and **20** as compared to
696 negative (untreated, DMSO and 1 μ g/mL INH) and positive (100 μ M metronidazole)
697 controls after 7 days of compound exposure. (C) Efficacy of compounds **1**, **9**, **10** and **16**

698 during growth of *Mtb* in J774 macrophages as compared to the negative (DMSO) and
699 positive (0.5 and 5 μ g/mL RIF) controls. Black horizontal bar indicates inoculum (day 0).

700 **FIG 4** The pyrimidinetrione amide lacks in vivo efficacy. (A) PK profile of compound **10**
701 after twice daily administration at 30 mg/kg. (B) *Mtb* bacterial burdens in infected mice
702 after 2 and 4 weeks of once daily or twice daily treatment at 30 and 15 mg/kg,
703 respectively. Treatment was initiated at 2 weeks after aerosol infection of the mice.

